

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Application of	Yoshiya Oda <i>et al.</i>	Confirmation No.	5776
Serial No.:	10/579,780	Art Unit:	1797
Filed:	May 18, 2006	Examiner:	Yelena G. Gakh
For:	<b>Quantitation Method Using Isotope Labeled Internal Standard Substance, Analysis System For Executing The Quantitation Method, And Program For The Analysis</b>	Atty. Docket No:	042715-5019-US

LETTER

Applicants call Examiner's attention to the attached reference (Liang et al., Ionization enhancement in atmospheric pressure chemical ionization and suppression in electrospray ionization between target drugs and stable-isotope-labeled internal standards in quantitative liquid chromatography/tandem mass spectrometry, *Rapid Communications in Mass Spectrometry*, 2003, 17: 2815-2821), which is placed in the file for the record.

Respectfully submitted,

Date: June 8, 2010  
  
MORGAN, LEWIS & BOCKIUS LLP  
1111 Pennsylvania Avenue, N.W.  
Washington, D.C. 20004  
Tel. Dir. (202) 739-5602

/ Mark J. Sullivan /  
Mark J. Sullivan Reg. No. 54,478

# Ionization enhancement in atmospheric pressure chemical ionization and suppression in electrospray ionization between target drugs and stable-isotope-labeled internal standards in quantitative liquid chromatography/tandem mass spectrometry<sup>†</sup>

H. R. Liang\*, R. L. Foltz, M. Meng and P. Bennett

Tandem Labs, A Division of NWT Inc., Salt Lake City, UT 84124, USA

Received 14 July 2003; Revised 14 October 2003; Accepted 15 October 2003

The phenomena of ionization suppression in electrospray ionization (ESI) and enhancement in atmospheric pressure chemical ionization (APCI) were investigated in selected-ion monitoring and selected-reaction monitoring modes for nine drugs and their corresponding stable-isotope-labeled internal standards (IS). The results showed that all investigated target drugs and their co-eluting isotope-labeled IS suppress each other's ionization responses in ESI. The factors affecting the extent of suppression in ESI were investigated, including structures and concentrations of drugs, matrix effects, and flow rate. In contrast to the ESI results, APCI caused seven of the nine investigated target drugs and their co-eluting isotope-labeled IS to enhance each other's ionization responses. The mutual ionization suppression or enhancement between drugs and their isotope-labeled IS could possibly influence assay sensitivity, reproducibility, accuracy and linearity in quantitative liquid chromatography/mass spectrometry (LC/MS) and liquid chromatography/tandem mass spectrometry (LC/MS/MS). However, calibration curves were linear if an appropriate IS concentration was selected for a desired calibration range to keep the response factors constant.

Copyright © 2003 John Wiley & Sons, Ltd.

Stable-isotope-labeled analogs are commonly used as internal standards (IS) in quantitative gas chromatography/mass spectrometry (GC/MS) and liquid chromatography/mass spectrometry (LC/MS).<sup>1–5</sup> The stable-isotope-labeled analogs are chemically and structurally the same as their target drugs but differ in molecular mass. The primary reason for utilizing a stable-isotope-labeled analog is to normalize the response of a given target drug to the response of its isotopic analog and thus compensate for variations in injection, sample preparation, instrumental parameters and matrix effects.<sup>3–13</sup> Previous reports have noted that, when constructing calibration curves with electrospray ionization (ESI), the peak areas of the co-eluting labeled IS decreases with increasing drug concentrations.<sup>12,13</sup> There are also some reports on ionization suppression in atmospheric pressure chemical ionization (APCI).<sup>14–16</sup> We have also noticed the above behavior when using ESI,<sup>16</sup> but with APCI we found that the peak

areas of stable-isotope-labeled IS increased when drug concentrations increased.

To further explore these phenomena and their possible influence on assay sensitivity, reproducibility, accuracy and linearity, we investigated the effects of ionization suppression in ESI and enhancement in APCI between nine drugs and their corresponding stable-isotope-labeled internal standards.

## EXPERIMENTAL

(R)- and (S)-Methadone and methadone-D<sub>2</sub> were purchased from Cerilliant (Austin, TX, USA). Fructose, sorbitol, fructose-<sup>13</sup>C<sub>6</sub>, sorbitol-<sup>13</sup>C<sub>6</sub>, fructose-D<sub>2</sub> and sorbitol-D<sub>2</sub> were purchased from Omicron Biochemicals, Inc. The other six drugs and their corresponding isotope-labeled internal standards are proprietary.

All chemicals were of analytical-reagent grade: hexane, methanol, isopropyl alcohol (IPA) and acetonitrile from EM Science (Gibbstown, NJ, USA); ammonium acetate and sodium bicarbonate from VWR Scientific Products (West Chester, PA, USA).

Chromatography was performed using a Shimadzu SCL-10A controller with CL-10AD pump and CTO-10A column

\*Correspondence to: H. R. Liang, Tandem Labs, A Division of NWT Inc., 1123 East 3900 South, Salt Lake City, UT 84124, USA. E-mail: harui@tandem-labs.com

<sup>†</sup>A part of this manuscript was presented as a poster at the 51st ASMS Conference on Mass Spectrometry and Allied Topics, Montreal, Canada, June 8–12, 2003.

oven (Shimadzu Scientific Instruments, Inc., Columbia, MD, USA). The autosampler was a PE series 200 (Perkin Elmer, Hopkinton, MA, USA). The API 3000 and 4000 tandem mass spectrometers were from MDS Sciex (Concord, Ontario, Canada).

The target drugs II–VII and their labeled IS were chromatographed using a Synergi Hydro-RP or Prodigy phenyl-3 column (30 × 2 mm, 5 µm particle size) with a mobile phase containing MeCN/5 mM ammonium acetate or 0.1% formic acid at a flow rate of 300 µL/min and an oven temperature of 40°C. (R)- and (S)-Methadone were chromatographed using a Chiral-AGP column (50 × 2 mm, 5 µm particle size) with a mobile phase containing isopropyl alcohol in ammonium acetate at a flow rate of 400 µL/min and at an oven temperature of 45°C. Fructose, sorbitol, fructose-<sup>13</sup>C<sub>6</sub>, sorbitol-<sup>13</sup>C<sub>6</sub>, fructose-D<sub>2</sub> and sorbitol-D<sub>2</sub> were chromatographed on a CapCell Pak 5 µ NH<sub>2</sub> column (UG-80A, 150 × 4.6 mm) with a mobile phase containing MeCN/MeOH/water.

The API 3000 and 4000 triple-quadrupole mass spectrometers were operated in either selected-ion monitoring (SIM) or selected-reaction monitoring (SRM) mode under optimized conditions for detection of positive or negative ions of drugs and IS formed by TurboIonSpray™ ionization (ESI) or heated nebulizer ionization (APCI). Drugs VIII and IX were studied using the API 4000, and the other drugs using the API 3000. Drugs V, VIII and IX were studied in negative ion mode and other drugs in positive ion mode.

The isotope-labeled IS and their target drugs were prepared in mobile phase. To explore the effects of matrix on the extent of suppression in ESI, 5000 ng/mL of fructose-<sup>13</sup>C<sub>6</sub>, 2500 ng/mL of sorbitol-<sup>13</sup>C<sub>6</sub>, fructose-D<sub>2</sub> and sorbitol-D<sub>2</sub> were spiked into 200 µL of human red blood cells and water in 10 replicates, and extracted with protein precipitation using acetonitrile.

The effect of flow rate on the extent of response suppression was studied by the injection of 20 µL of 1000 ng/mL drug II and the post-column infusion of drug II-13C<sub>6</sub> at the rate of 20 µL/min while changing the chromatographic flow rate from 0.1 to 0.8 mL/min.

The experiments to investigate ionization suppression and enhancement were performed using a post-column infusion

system in which a constant flow of a drug (or its IS) was infused post-column into the MS detector and the IS (or the drug) injected by an autosampler onto the analytical column.<sup>17,18</sup> The purpose of the post-column infusion with a drug (or its IS) is to raise the background level so that the suppression by the IS (or the drug) will show as negative peaks and the enhancement will show as positive peaks. The extent of ionization suppression or enhancement was also measured by comparing the peak areas of drug or IS from solutions containing only drug or IS or both of them.

The chromatographic peaks for the target drugs and their corresponding stable-isotope-labeled IS were integrated using Analyst software (version 1.2) with a smoothing factor of one. Quantitation was based on linear regression analysis of calibration curves (weighted 1/x) using the analyte to IS area ratio vs. target concentration utilizing Watson® DMUIMS software (version 6.1.1.04).

## RESULTS AND DISCUSSION

Table 1 lists information on the nine investigated drugs. These included basic, acidic and neutral compounds, and their deuterium- or <sup>13</sup>C-labeled internal standards.

### Ionization suppression in ESI

#### Ionization suppression between target drugs and their stable isotope-labeled IS in SIM and SRM modes

The results showed that, when using ESI, all target drugs suppressed the ionization responses of their co-eluting labeled IS in both SIM and SRM modes, and likewise the labeled IS suppressed the ionization responses of the corresponding target drugs. Figure 1 shows a representative example of the results. This suppression can be understood in terms of Enke's model of ESI ion generation,<sup>19,20</sup> which involves a result of the competition among ions for the limited number of excess charge sites on the generated droplet during ESI. This model predicts the response curves of singly charged ionic analytes as a function of the concentration of electrolyte and other analytes. The extents of suppression of signals from the nine target drugs by their corresponding stable-isotope-labeled IS are listed in Table 1.

Table 1. Nine investigated target drugs and their corresponding isotope-labeled internal standards (IS)

Analytes	Analytes labeled IS					
	[M+H] <sup>+</sup> (m/z)	IS	[M+H] <sup>+</sup> (m/z)	Isotopic contribution*	Structure type	Extent of suppression** %
I (R, S)-methadone	310	D <sub>2</sub>	313	0.2%	Tertiary amine	72
II	409	<sup>13</sup> C <sub>6</sub>	413	<0.2%	Primary amine	88
III	478	<sup>13</sup> C <sub>2</sub> , D <sub>2</sub>	482	<0.2%	Secondary amine	84
IV	506	<sup>13</sup> C <sub>2</sub> , D <sub>2</sub>	510	<0.2%	Tertiary amine	82
V	274	D <sub>4</sub>	478	0.2%	Acidic	80
VI	249	D <sub>3</sub>	253	0.5%	Neutral	62
VII	265	D <sub>3</sub>	268	0.5%	Neutral	64
VIII sorbitol	181	<sup>13</sup> C <sub>6</sub>	187	<0.2%	Neutral	38
IX fructose	179	<sup>13</sup> C <sub>6</sub>	185	<0.2%	Neutral	40

\* Isotopic contribution indicates contribution of naturally occurring isotopic abundance of drugs to their corresponding isotope-labeled IS response assuming equal concentrations of the drug and the IS.

\*\* Extent of suppression was calculated as 100 × [(Intensity of signal from drug by post-column infusion – signal intensity of drug suppressed due to the on-column injection of their IS)/intensity of signal from drug by post-column infusion]. Concentrations of drugs and their IS: 10 µg/mL; injection volume: 20 µL; infusion rate: 20 µL/min.

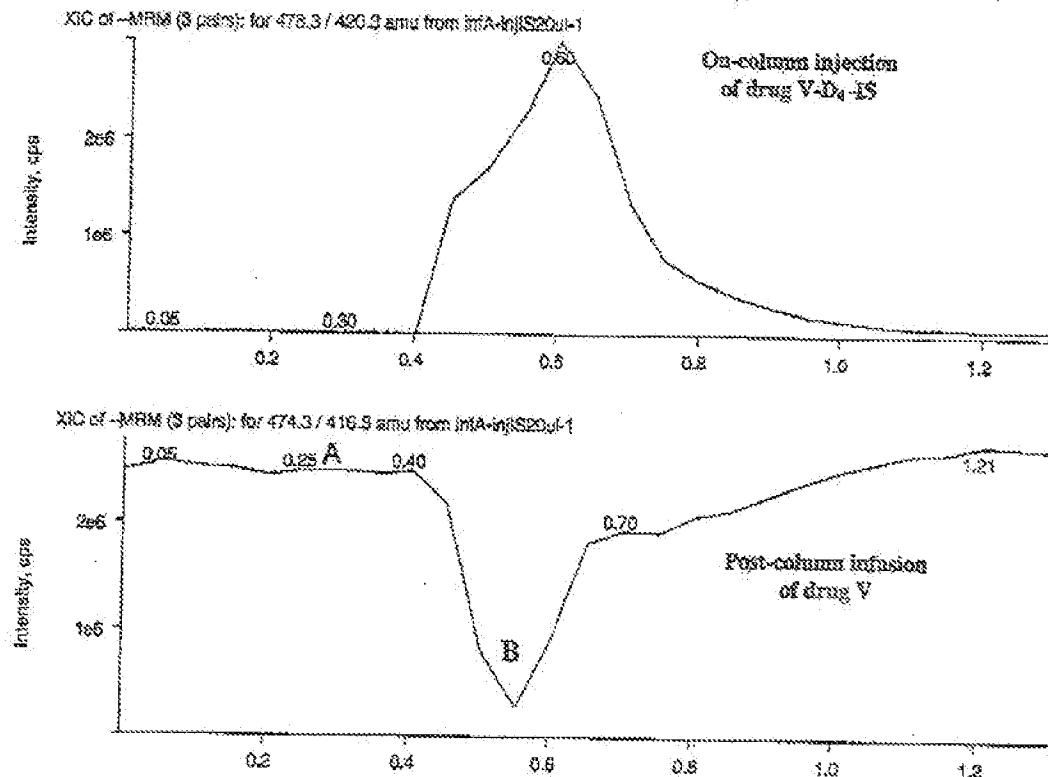


Figure 1. ESI response of drug V suppressed ~80% by its  $D_3$ -IS during the elution window of  $D_4$ -IS. (A) Intensity of  $D_0$  by post-column infusion and (B) intensity of  $D_0$  suppressed due to the on-column injection of  $D_3$ . Using a post-column infusion system, drug V was infused post-column and its  $D_3$  analog was injected on-column. For other conditions, see Experimental section. Concentrations of drug V and its  $D_3$  analog were 10  $\mu$ g/mL, injection volume: 20  $\mu$ L; infusion rate: 20  $\mu$ L/min. The extent of suppression % was calculated as:  $100 \times \{(A - B)/A\}$ .

#### Factors affecting the extent of suppression

**Concentrations of investigated drugs and IS.** Figure 2 illustrates the mutual ionization suppression between methadone

(from 50–10000 ng/mL) and its  $D_3$ -IS (25–5000 ng/mL). The results demonstrate that, as the concentration of the  $D_3$ -IS increased, the greater the suppression of methadone  $D_0$  was caused by its  $D_3$ -IS, and vice versa. For example, at the concentrations 50/25 ng/mL of  $D_0$ / $D_3$ , the suppression was approximately 10%/30%, respectively. At the 5000/2500 ng/mL concentration level the suppression was 25%/50%. The extent of suppression in each drug-IS pair was concentration-dependent in a nonlinear fashion.

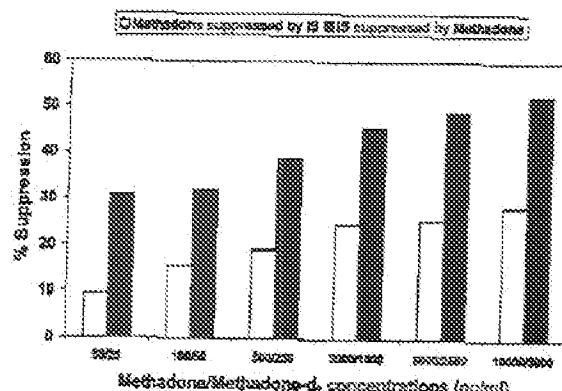


Figure 2. ESI suppression between methadone and methadone- $D_3$ -IS. The white bars indicate the extent of suppression of  $D_0$  ionization by  $D_3$  by comparing the LC/MS/MS peak areas of  $D_0$  from the solutions containing only  $D_0$  with those containing both  $D_0$  and  $D_3$ . The black bars indicate the extent of suppression of  $D_3$  ionization by  $D_0$  by comparing the peak areas of  $D_3$  from the solutions containing only  $D_3$  with those containing both of them. Concentrations of  $D_0$ : 50–10000 ng/mL; concentrations of  $D_3$ : 25–5000 ng/mL. For other conditions, see Experimental section.

**Hydrophobicity of investigated drugs.** The extent of suppression was related to the structures of the investigated drugs (Table 1). Under optimized LC/MS/MS conditions, the extent of suppression for each investigated drug by its IS was different when the concentration, injection volume, and infusion rate of drugs or IS were exactly the same. The hydrophobicity was determined based on organic vs. aqueous partitioning. Generally, the level of suppression was correlated with the hydrophobicity of the compounds; the more hydrophobic the compound, the lower the level of suppression. However, fructose and sorbitol, both hydrophilic compounds, did not comply with this general finding. Only 35–40% of sorbitol or fructose response was suppressed while ~60% of the signals of two hydrophobic drugs, VI and VII, were suppressed.

**Matrix effects.** Fructose-<sup>13</sup>C<sub>6</sub> and sorbitol-<sup>13</sup>C<sub>6</sub> and/or fructose- $D_2$  and sorbitol- $D_2$  were spiked into human red blood

**Table 2.** Ionization suppression in ESI between fructose-<sup>13</sup>C<sub>6</sub> (sorbitol-<sup>13</sup>C<sub>2</sub>) and fructose-D<sub>2</sub> (sorbitol-D<sub>2</sub>) extracted from human red blood cells (RBCs) and water, respectively

Peak areas of fructose-, sorbitol- <sup>13</sup> C <sub>6</sub> and D <sub>2</sub> spiked only with <sup>13</sup> C <sub>6</sub> or D <sub>2</sub> in water				
(n = 10)	fructose- <sup>13</sup> C <sub>6</sub>	fructose-D <sub>2</sub>	sorbitol- <sup>13</sup> C <sub>6</sub>	sorbitol-D <sub>2</sub>
AVE	786.09000	330.10000	258.62500	311.45600
Std Dev	377.585	110.6300	71.8470	144.6033
CV%	0.48	3.3	2.7	4.6
Peak areas of fructose-, sorbitol- <sup>13</sup> C <sub>6</sub> and D <sub>2</sub> spiked with both <sup>13</sup> C <sub>6</sub> and D <sub>2</sub> in water				
(n = 10)	fructose- <sup>13</sup> C <sub>6</sub>	fructose-D <sub>2</sub>	sorbitol- <sup>13</sup> C <sub>6</sub>	sorbitol-D <sub>2</sub>
Ave	667.36000	218.33000	224.85000	278.81500
Std Dev	146.3711	55.5785	27.4357	184.554
CV%	2.1	2.5	1.2	6.6
Suppression %	15.1	33.8	15.0	10.4
Peak areas of fructose-, sorbitol- <sup>13</sup> C <sub>6</sub> and D <sub>2</sub> spiked only with <sup>13</sup> C <sub>6</sub> or D <sub>2</sub> in human RBCs				
(n = 10)	fructose- <sup>13</sup> C <sub>6</sub>	fructose-D <sub>2</sub>	sorbitol- <sup>13</sup> C <sub>6</sub>	sorbitol-D <sub>2</sub>
AVE	489.84500	608.6050	165.76500	834.0900
Std Dev	109.15607	244.6660	49.33484	267.5833
CV%	22.2	40.2	29.7	32.0
Peak areas of fructose-, sorbitol- <sup>13</sup> C <sub>6</sub> and D <sub>2</sub> spiked with both <sup>13</sup> C <sub>6</sub> and D <sub>2</sub> in human RBCs				
(n = 10)	fructose- <sup>13</sup> C <sub>6</sub>	fructose-D <sub>2</sub>	sorbitol- <sup>13</sup> C <sub>6</sub>	sorbitol-D <sub>2</sub>
Ave	131.68500	297.9050	443.5550	493.3700
Std Dev	150.5430	362.250	414.293	391.595
CV%	11.4	12.1	9.340304	7.937161
Suppression %	73.1	51.0	73.2	40.8

The suppression (%) of fructose- and sorbitol-<sup>13</sup>C<sub>6</sub> by their D<sub>2</sub> (or D<sub>2</sub> by <sup>13</sup>C<sub>6</sub>) was determined by comparing the peak areas of <sup>13</sup>C<sub>6</sub> (or D<sub>2</sub>) from solutions containing only <sup>13</sup>C<sub>6</sub> (or D<sub>2</sub>), or both <sup>13</sup>C<sub>6</sub> and D<sub>2</sub>. Concentration of fructose-<sup>13</sup>C<sub>6</sub>: 5000 ng/mL; concentration of sorbitol-<sup>13</sup>C<sub>6</sub> or D<sub>2</sub>: 2500 ng/mL; injection volume: 10  $\mu$ L.

cells and water, respectively, and extracted with the exact same procedures. As shown in Table 2 there were significant matrix effects; there were significant differences in signal suppression for fructose-<sup>13</sup>C<sub>6</sub> and sorbitol-<sup>13</sup>C<sub>6</sub> by their D<sub>2</sub> (or D<sub>2</sub> by <sup>13</sup>C<sub>6</sub>) in human red blood cells compared with water. The signals were suppressed more in human red blood cells compared with those in water. This result can also be understood in terms of Enke's model of ESI ion generation,<sup>19,20</sup> a result of the competition among ions for the limited number of excess charge sites on the generated droplet during ESI.

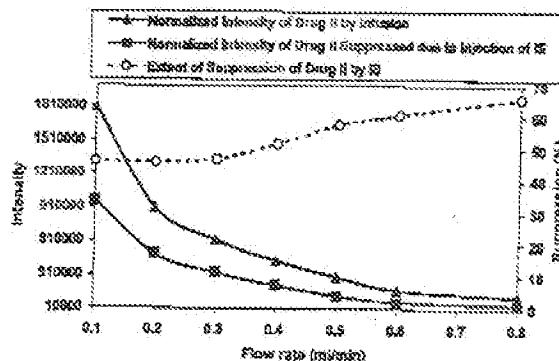
**Flow rate.** Generally, the higher the flow rate, the higher the level of suppression. Figure 3 shows a representative result. The signal suppression of analytes tended to be lower at lower flow rates. This can also be explained by Enke's model in which the sum of the excess charge concentration at the surface is inversely proportional to flow rate.<sup>19,20</sup>

#### Effect of ionization suppression on assay sensitivity, reproducibility, accuracy and linearity

Table 3 shows the influence of (R)-methadone-D<sub>3</sub> (IS) concentrations on the sensitivity, reproducibility, accuracy and linearity for (R)-methadone. The IS concentrations influenced the detection limit and the lower limit of quantitation (LLOQ) of the assay. As an example of this effect, without addition of the IS the response for 1 ng/mL of (R)-methadone was observed as a peak with signal-to-noise (S/N) ratio > 10, but in the presence of 200 ng/mL of IS the response for 1 ng/mL of (R)-methadone was totally suppressed by the IS. When the IS concentration was increased to 10 000 ng/mL, the responses for up to 10 ng/mL of (R)-methadone were totally suppressed and the response of 25 ng/mL of (R)-methadone was suppressed up to 93%.

The IS concentration also influenced the reproducibility, response factors, accuracy and linearity of the assay because

the mutual suppression was concentration-dependent. The results appeared to be contrary to those of a previous report,<sup>12</sup> in which Sojo *et al.* found that the mutual suppression did not have an effect on the calibration curve and quantitation of target drugs. Based on the present results, the reproducibility was better for the IS responses and worse for the response factors and accuracy of the unlabeled drug; also the linear range was narrower with different slope and correlation of determination in the calibration series in the presence of 10 000 ng/mL of IS than in the presence of 200 ng/mL of IS. In the presence of 10 000 ng/mL of IS, the



**Figure 3.** Effect of flow rate on suppression of ESI of drug II by drug II-<sup>13</sup>C<sub>6</sub>-IS. (A) Intensity of drug II by post-column infusion normalized to the flow rate and (B) intensity of drug II suppressed due to the on-column injection of <sup>13</sup>C<sub>6</sub> normalized to the flow rate. The extent of suppression % was calculated as:  $100 \times ((A - B)/A)$ . Using a post-column infusion system, drug II was infused post-column and its <sup>13</sup>C<sub>6</sub> was injected on-column. Concentrations of drug II and its <sup>13</sup>C<sub>6</sub>: 1  $\mu$ g/mL; injection volume: 20  $\mu$ L; infusion rate: 20  $\mu$ L/min. For other conditions, see Experimental section.

Table 3. Influence of (R)-methadone-D<sub>3</sub> internal standard (IS) concentrations on sensitivity, precision, accuracy and linearity of (R)-methadone in calibration curves (n = 4)

Concentration of (R)-methadone (ng/mL)	IS peak area	Response factor*	Accuracy	(R)-Methadone peak area	Suppression %**
<i>Without addition of IS</i>					
1	2001	152.0	100.1	3001.1	
5	2486	163.2	12.028		
25	2573	230.9	64.331		
50	1957	180.5	97.827		
100	1830	173.5	182.960		
500	1344	133.4	67.830		
2000	966	104.6	198.1300		
5000	627	75.7	263.9700		
8500	543	85.4	429.5100		
10000	527	N/A	527.4000		
AVE	1295	140.6			
STD DEV	763	72.4			
CV%	56.9	51.5			
<i>With addition of 200 ng/mL of IS</i>					
1	192.880	0.00000	N/A	N/A	N/A
5	180.610	0.01079	112.1	5812	26.7
25	165.640	0.01086	92.5	52.365	18.6
50	165.520	0.01068	68.5	96.432	1.4
100	163.370	0.01059	86.6	175.360	4.2
500	113.550	0.01107	89.4	628.500	6.4
2000	71.356	0.01286	103.7	183.5300	5.0
5000	55.176	0.01255	101.1	249.8700	5.1
8500	41.192	0.01310	105.6	458.8000	0.2
10000	39.827	0.01175	94.7	524.4400	0.6
AVE	101.963	0.01166	95.3		
STD DEV	56.921	0.00103	7.3		
CV%	55.8	8.8	7.7		
R <sup>2***</sup>	0.999 from 5 to 10 000 ng/mL				
<i>With addition of 10 000 ng/mL of IS</i>					
1	257.4600	N/A	N/A	N/A	N/A
5	214.3500	N/A	N/A	N/A	N/A
25	211.9600	0.00007	181.0	4473	93.0
50	224.9300	0.00012	122.1	13.423	86.3
100	218.8100	0.00016	102.4	35.171	80.8
500	229.9800	0.00021	93.7	271.950	59.5
2000	193.0100	0.00024	99.1	109.3200	43.4
5000	148.6000	0.00023	96.4	173.7500	34.0
5500	170.4800	0.00025	102.7	362.1000	21.2
10000	226.1100	0.00019	89.0	419.0100	20.6
AVE	201.9850	0.00019	112.0		
STD DEV	289.726	0.00006	29.2		
CV%	14.3	34.9	26.0		
R <sup>2***</sup>	0.954 from 25 to 10 000 ng/mL				

\* Response factor: response factor refers to the peak area ratio (drug/internal standard) vs. drug concentration.

\*\* Suppression %: %[(R)-methadone peak areas obtained without addition of IS] - [(R)-methadone peak areas obtained with addition of IS)] / [(R)-methadone obtained without addition of IS].

\*\*\* R<sup>2</sup>: correlation of determination.

responses of (R)-methadone at low concentrations were suppressed to a greater extent (up to 93%) than at higher concentrations (20%). In the presence of 200 ng/mL of IS, the extent of suppression of the drug responses varied from 26 to 0.3%.

Therefore, it is important to select an appropriate IS concentration for a desired calibration range to keep calibration curves linear. Generally, IS concentration should not be too high. An appropriate IS concentration depends on the investigated drugs and other experimental parameters and thus should be determined by experiment. For the methadone assay, the appropriate IS concentration was

found to be 200 ng/mL for the desired linear range from 5–10 000 ng/mL, with R<sup>2</sup> = 0.999 and overall %CV of response factors = 8.8%.

On the other hand, the extent of suppression of the IS responses depended on the drug concentrations. The suppression of the IS response by its target drug caused the apparent poor reproducibility of the IS response. Figure 4 shows that the peak areas of co-eluting labeled (R)- and (S)-methadone-D<sub>3</sub> decreased as the concentration of (R)- and (S)-methadone increased in the calibration series. This behavior was also IS concentration-dependent and was not significant in the presence of 10 000 ng/mL of IS (Table 3).

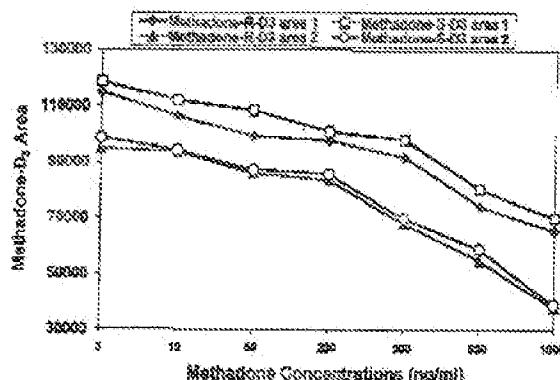


Figure 4. Electrospray peak areas of co-eluting labeled (R)- and (S)-methadone-D<sub>4</sub>-IS decreasing with increasing (R)- and (S)-methadone concentrations in a calibration series. Areas 1 and 2 are from results obtained on different days. For conditions, see Experimental section.

#### Ionization enhancement in APCI

##### Ionization enhancement between target drugs and their stable-isotope-labeled IS in SIM and SRM modes

In contrast to the ESI results, seven out of the nine investigated target drugs enhanced the ionization responses of the

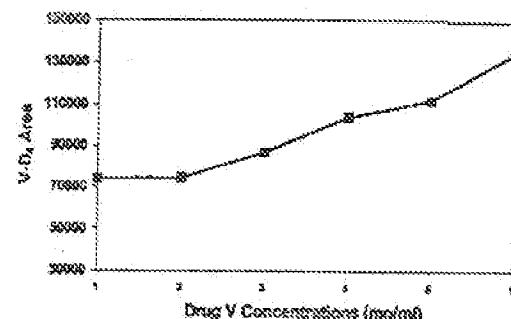


Figure 6. LC/APCI-MS/MS peak areas of D<sub>4</sub>-IS increase with increasing drug V concentration. For conditions, see Experimental section.

corresponding labeled IS in both SIM and SRM modes, and likewise the labeled IS enhanced the responses of the target drugs with APCI. The responses of drugs I-VII were enhanced 2–7 times by their corresponding IS. Figure 5 shows that the response of drug V was enhanced ~7 times by its D<sub>4</sub> analog. However, the ionization responses of 10 µg/mL of fructose and sorbitol by post-column infusion were not enhanced by on-column injection of 10 µg/mL of fructose-<sup>13</sup>C<sub>6</sub> or sorbitol-<sup>13</sup>C<sub>6</sub>, but rather were suppressed

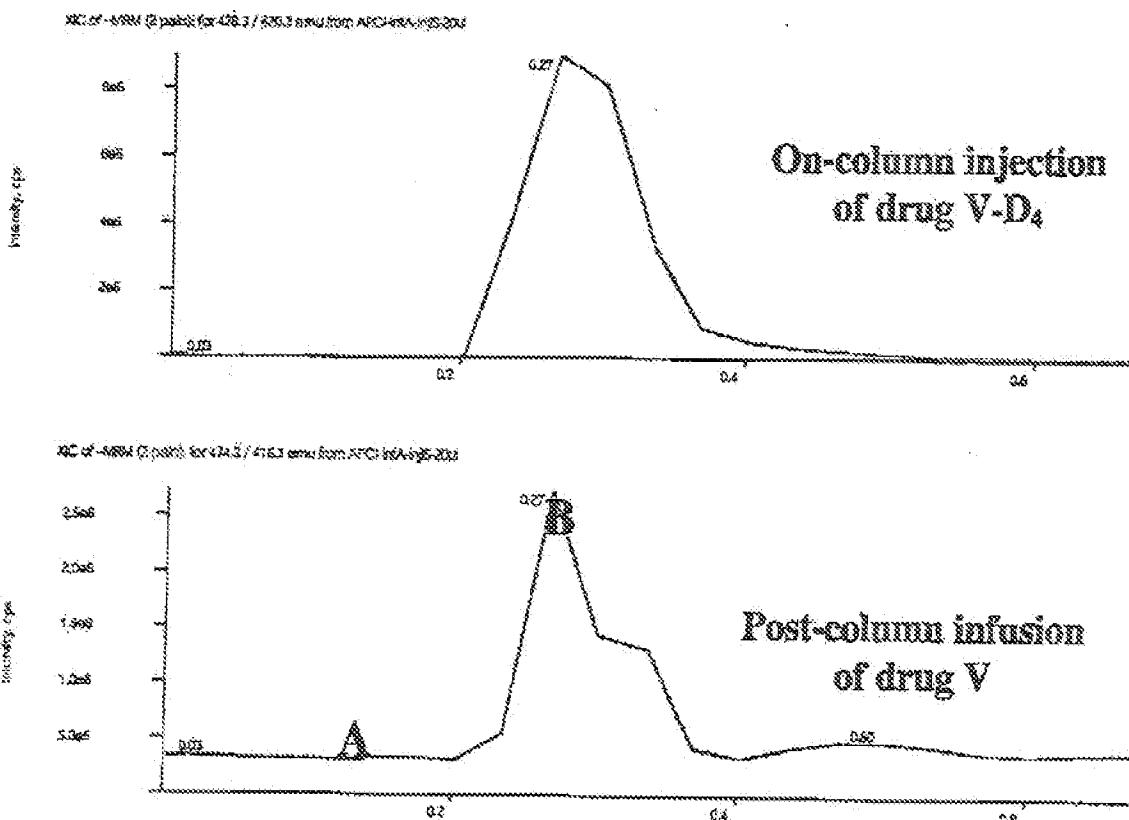


Figure 5. APCI response of drug V enhanced ~7 times by its D<sub>4</sub>-IS during the retention time window of the D<sub>4</sub>-IS. (A) Intensity of D<sub>9</sub> by post-column infusion and (B) Intensity of D<sub>9</sub> enhanced due to the on-column injection of D<sub>4</sub>. Using a post-column infusion system, drug V was infused post-column and its D<sub>4</sub>-IS was injected on-column. The extent of enhancement was calculated as  $(B - A)/A$ . Concentrations of drug V and its D<sub>4</sub>-IS: 10 µg/mL; injection volume: 20 µL; infusion rate: 20 µL/min. For other conditions, see Experimental section.

**Table 4.** Precision of IS peak area (V-D<sub>4</sub>), response factor and accuracy of drug V in calibration curves (n=2)

Concentration of drug V (mg/mL)	IS peak area	Response factor	Accuracy
0.400	74056	2.81	99
0.800	76431	3.03	105
2.00	86968	3.03	104
4.00	103970	2.87	98
6.00	112240	2.76	98
7.20	134690	3.07	105
AVE	97725	2.93	101
STDV	23810	0.13	43
CV%	24.3	4.4	43

by fructose-<sup>13</sup>C<sub>6</sub> or sorbitol-<sup>13</sup>C<sub>6</sub> when their concentrations were increased to 50 µg/mL.<sup>14,15</sup> Explanations for these results are still being explored.

#### Extent of enhancement and concentrations of investigated drugs

In some cases, the enhancement of IS signals by their corresponding drugs was concentration-dependent, and the peak areas for the IS increased with increasing drug concentrations in calibration curves. Figure 6 shows that the peak areas of the co-eluting V-D<sub>4</sub> increased with increasing drug V concentrations. However, this behavior was not significant for five of the nine investigated target drug-IS pairs.

#### Effect of ionization enhancement on assay sensitivity, reproducibility, accuracy and linearity

Ionization enhancement of drugs by their stable-isotope-labeled IS in APCI could possibly improve the detection limits and the LLOQ of the assay for some drugs. The increase of IS responses with increasing drug concentrations in the calibration series resulted in poor apparent reproducibility for the IS signal. However, calibration curves were linear if an appropriate internal standard concentration was selected for a desired calibration range to keep the response factor constant (Table 4).

#### Effect of natural isotopic contribution from drugs on enhancement of IS

The contributions of naturally occurring isotopic abundance of drugs to the signals for their corresponding isotope-labeled IS are shown in Table 1. The degree of enhancement of isotope-labeled IS was substantially greater than could be accounted for by the naturally occurring isotopic abundances of their target drugs.

#### Effect of purity of IS on enhancement of drugs

In the present study, the isotopic purities of all isotope-labeled IS used were over 99%. Therefore, the presence of any significant D<sub>4</sub>-IS impurities was ruled out.

## CONCLUSIONS

Ionization enhancement under APCI conditions between target drugs and co-eluting isotope-labeled IS was investigated in quantitative LC/MS and LC/MS/MS for the first time. In APCI, seven out of nine investigated target drugs and their co-eluting isotope-labeled IS were found to enhance each other's ionization responses. In ESI, all investigated target drugs and their co-eluting isotope-labeled IS were found to suppress each other's ionization responses. The mutual ionization enhancement and suppression between drugs and their isotope-labeled IS can influence assay sensitivity, reproducibility, accuracy and linearity. Linear calibration curves can be maintained if an appropriate IS concentration is selected for a desired calibration range to keep the response factor constant.

## REFERENCES

- Edlund O, Bowers L, Henion J, Covey TR. *J. Chromatogr.* 1989; 497: 49.
- Jaruga P, Birincioğlu M, Rodriguez H, Dizdaroglu M. *Biochemistry* 2002; 41: 3703.
- Chang WT, Smith J, Liu RH. *J. Forensic Sci.* 2002; 47: 873.
- Liu RH, Lin D, Chang W, Liu C, Tsay W, Li J, Kuo T. *Anal. Chem.* 2002; 619: 27A.
- Ellerbe P, Meldelein S, Sniegosky LT, Welch MJ, White E. *Anal. Chem.* 1989; 61: 1710.
- Jacob P, Wilson M, Yu L, Mendelson J, Jones ST. *Anal. Chem.* 2002; 74: 5290.
- Vinsal F, Fogliano V, Schieberle P, Hofmann T. *J. Agric. Food Chem.* 1999; 47: 5084.
- Kuklenyik Z, Ashley DL, Calafat AM. *Anal. Chem.* 2002; 74: 2053.
- Magni P, Pereira S, Leoni M, Criscenti G, Kienle GM. *J. Mass Spectrom.* 2001; 36: 670.
- Dalluge J, Hashizume T, McCloskey JA. *Nucleic Acids Res.* 1996; 24: 3242.
- Brown FR, Draper WM. *Biol. Mass Spectrom.* 1991; 20: 515.
- Soto LE, Lum G, Clase P. *Analyst* 2003; 128: 51.
- Zhou W, Zhou S, Tollefson KL, Jiang X, Weng N. *Proc. 51st ASMS Conf. Mass Spectrometry and Allied Topics*, Montreal, Canada, June 8–12, 2003.
- Mei H, Hsieh Y, Nardo C, Xu X, Wang S, Ng K, Korfner W. *Rapid Commun. Mass Spectrom.* 2003; 17: 97.
- Hout MWJV, Niederlander HAG, Zeeuw Rada, Jong GJde. *Rapid Commun. Mass Spectrom.* 2003; 17: 245.
- Liang HR, Folz RL, Meng M, Bennett P. *Proc. 51st ASMS Conf. Mass Spectrometry and Allied Topics*, Montreal, Canada, June 8–12, 2003.
- Miller-Stein C, Bonfiglio R, Olah TA, King RC. *Am. Pharm. Rev.* 2000; 54.
- King RC, Bonfiglio R, Fernandez-Metzler C, Miller-Stein C, Olah TA. *J. Am. Soc. Mass Spectrom.* 2000; 11: 942.
- Enke CG. *Anal. Chem.* 1997; 69: 4835.
- Cech NB, Enke CG. *Anal. Chem.* 2001; 73: 4632.